Express Mail EV 330856466 US Mailed August 4, 2003

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of:

HARRIS et al.

Application No.: Not Yet Assigned

Filed: August 4, 2003

For: SCREENING ASSAYS FOR COMPOUNDS THAT CAUSE APOPTOSIS

Customer No.: 20350

Mail Stop Patent Application Commissioner for Patents P.O. Box 1450 Alexandria, VA 22313-1450

Sir:

Confirmation No.: Not Yet Assigned

Examiner:

Not Yet Assigned

Technology Center/Art Unit: Not Yet

Assigned

REQUEST TO ACCEPT IN THE PRESENT CASE THE SEQUENCE LISTING FILED IN THE PARENT CASE

Applicants have previously filed on October 28, 1996 a sequence listing and accompanying computer readable diskette in the parent Application No. 08/359,316. Applicants request the use of the previously filed diskette containing the sequence information in computer readable form in the parent application in lieu of providing a diskette herewith. The diskette submitted to the USPTO in the parent application on October 28, 1996 is the only computer-readable form filed. The parent application information data is as follows:

Applicants: Harris et al. Serial No. 08/359,316 Filed: December 19, 1994

Title: SCREENING ASSAYS FOR COMPOUNDS THAT CAUSE APOPTOSIS

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As required by 1.821(f), applicants state that the content of the above-identified computer readable diskette submitted in the parent application is the same as the content of the hard copy of the Sequence Listing (pages 37-43) submitted to the USPTO in the parent application and herewith in the accompanying divisional application.

Respectfully submitted,

Kenneth A. Weber Reg. No. 31,677

TOWNSEND and TOWNSEND and CREW LLP Two Embarcadero Center, 8th Floor San Francisco, California 94111-3834

Tel: 415-576-0200 Fax: 415-576-0300 Attachments KAW:dk

KAW:dk 60010106 v1

Transmittal Sheet

Atty. Docket No. 15280-225000

addressed to:

I hereby certify that this is being deposited with the United

States Postal Service as first class mail in an envelope

Assistant Commissioner for Patents

Sterry Barton

Washington, D. C. 20231.

Date: October 28, 1996

Date October 28, 1996

TOWNSEND and TOWNSEND and CREW LLP

Two Embarcadero Center Eight Floor

San Francisco, CA 94111-3834

(415) 576-0200

(415) 576-0300 FAX

In re application of:

Curtis C. Harris et al.

Serial No.: 08/359,316

Filed: December 19, 1994

Group Art Unit: 1811

For:

SCREENING ASSAYS FOR

COMPOUNDS THAT CAUSE

APOPTOSIS

THE ASSISTANT COMMISSIONER FOR PATENTS

Washington, D.C. 20231

Sir:

Transmitted herewith for the above-identified application are:

[X] Petition to Extend Time [CFR § 1.136(a)] with fee authorization \$110.00

- [X] Declaration with Exhibits 1-4 [CFR § 1.131]
- [X] Amendment [37 CFR § 1.115]
- [X] Substitute Paper Copy and Computer Readable Form of Sequence Listing [37 C.F.R. §§ 1.821-1.825]
- [X] Fee

[] No fee is due.

Please charge Deposit Account No. 20-1430 as follows:

[X] Any additional fees associated with this paper or during the pendency of this application.

Two copies of this sheet are enclosed.

TOWNSEND and TOWNSEND and CREW LLP

-trall

Kenneth A. Weber Reg. No.: 31,677

Attorneys for Applicants

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: HARRIS, Curtis C. WANG, Xin Wei HOEIJMAKERS, Jan H.J.
- (ii) TITLE OF INVENTION: SCREENING ASSAYS FOR COMPOUNDS THAT CAUSE APOPTOSIS
- (iii) NUMBER OF SEQUENCES: 6
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Townsend and Townsend Khourie and Crew
 - (B) STREET: Steuart Street Tower, One Market Plaza
 - (C) CITY: San Francisco
 - (D) STATE: California
 - (E) COUNTRY: US
 - (F) ZIP: 94105-1493
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- - (B) FILING DATE: 19-DEC-1994
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Kruse, Norman J.
 - (B) REGISTRATION NUMBER: 35,235
 - (C) REFERENCE/DOCKET NO: 15280-225000
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (415) 543-9600 (B) TELEFAX: (415) 543-5043

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 393 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
- (ix) FEATURE:
 - (A) NAME/KEY: Protein
 - (B) LOCATION: 1..393
 - (D) OTHER INFORMATION: /note= "Amino acid sequence of human wild-type p53."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
- Met Glu Glu Pro Gln Ser Asp Pro Ser Val Glu Pro Pro Leu Ser Gln
 1 5 10 15
- Glu Thr Phe Ser Asp Leu Trp Lys Leu Leu Pro Glu Asn Asn Val Leu 20 25 30
- Ser Pro Leu Pro Ser Gln Ala Met Asp Asp Leu Met Leu Ser Pro Asp 35 40 45
- Asp Ile Glu Gln Trp Phe Thr Glu Asp Pro Gly Pro Asp Glu Ala Pro 50 55 60
- Arg Met Pro Glu Ala Ala Pro Arg Val Ala Pro Gly Pro Ala Ala Pro 65 70 75 80
- Thr Pro Ala Ala Pro Ala Pro Ala Pro Ser Trp Pro Leu Ser Ser Ser 85 90 95
- Val Pro Ser Gln Lys Thr Tyr Gln Gly Ser Tyr Gly Phe Arg Leu Gly 100 105 110
- Phe Leu His Ser Gly Thr Ala Lys Ser Val Thr Cys Thr Tyr Ser Pro 115 120 125
- Ala Leu Asn Lys Met Phe Cys Gln Leu Ala Lys Thr Cys Pro Val Gln 130 135 140
- Leu Trp Val Asp Ser Thr Pro Pro Pro Gly Thr Arg Val Arg Ala Met 145 150 155 160
- Ala Ile Tyr Lys Gln Ser Gln His Met Thr Glu Val Val Arg Arg Cys 165 170 175
- Pro His His Glu Arg Cys Ser Asp Ser Asp Gly Leu Ala Pro Pro Gln 180 185 190
- His Leu Ile Arg Val Glu Gly Asn Leu Arg Val Glu Tyr Leu Asp Asp 195 200 205
- Arg Asn Thr Phe Arg His Ser Val Val Val Pro Tyr Glu Pro Pro Glu 210 215 220
- Val Gly Ser Asp Cys Thr Thr Ile His Tyr Asn Tyr Met Cys Asn Ser 225 230 240
- Ser Cys Met Gly Gly Met Asn Arg Arg Pro Ile Leu Thr Ile Ile Thr 245 250 255

- Leu Glu Asp Ser Ser Gly Asn Leu Leu Gly Arg Asn Ser Phe Glu Val
- Arg Val Cys Ala Cys Pro Gly Arg Asp Arg Arg Thr Glu Glu Glu Asn 280
- Leu Arg Lys Lys Gly Glu Pro His His Glu Leu Pro Pro Gly Ser Thr
- Lys Arg Ala Leu Pro Asn Asn Thr Ser Ser Pro Gln Pro Lys Lys 310
- Lys Pro Leu Asp Gly Glu Tyr Phe Thr Leu Gln Ile Arg Gly Arg Glu
- Arg Phe Glu Met Phe Arg Glu Leu Asn Glu Ala Leu Glu Leu Lys Asp
- Ala Gln Ala Gly Lys Glu Pro Gly Gly Ser Arg Ala His Ser Ser His
- Leu Lys Ser Lys Lys Gly Gln Ser Thr Ser Arg His Lys Lys Leu Met 375

Phe Lys Thr Glu Gly Pro Asp Ser Asp 385 390

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
- (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..19
 - (D) OTHER INFORMATION: /note= "Peptide # p53cp: amino acid sequence of p53 peptide."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Ser His Leu Lys Ser Lys Lys Gly Gly Ser Thr Ser Arg His Lys Lys

Leu Met Phe

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 781 amino acids

 - (B) TYPE: amino acid (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
- (ix) FEATURE:
 - (A) NAME/KEY: Protein
 - (B) LOCATION: 1..781
 - (D) OTHER INFORMATION: /note= "Amino acid sequence of human XPB."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
- Met Gly Lys Arg Asp Arg Ala Asp Arg Asp Lys Lys Lys Ser Arg Lys
- Arg His Tyr Glu Asp Glu Glu Asp Asp Glu Glu Asp Ala Pro Gly Asn
- Asp Pro Gln Glu Ala Val Pro Ser Ala Ala Gly Lys Gln Val Asp Glu
- Ser Gly Thr Lys Val Asp Glu Tyr Gly Ala Lys Asp Tyr Arg Leu Gln
- Met Pro Leu Lys Asp Asp His Thr Ser Arg Pro Leu Trp Val Ala Pro
- Asp Gly His Ile Phe Leu Glu Ala Phe Ser Pro Val Tyr Lys Tyr Ala
- Gln Asp Phe Leu Val Ala Ile Ala Glu Pro Val Cys Arg Pro Thr His 100
- Val His Glu Tyr Lys Leu Thr Ala Tyr Ser Leu Tyr Ala Ala Val Ser
- Val Gly Leu Gln Thr Ser Asp Ile Thr Glu Tyr Leu Arg Lys Leu Ser
- Lys Thr Gly Val Pro Asp Gly Ile Met Gln Phe Ile Lys Leu Cys Thr
- Val Ser Tyr Gly Lys Val Lys Leu Val Leu Lys His Asn Arg Tyr Phe 165 170 175
- Val Glu Ser Cys His Pro Asp Val Ile Gln His Leu Leu Gln Asp Pro
- Val Ile Arg Glu Cys Arg Leu Arg Asn Ser Glu Gly Glu Ala Thr Glu 200
- Leu Ile Thr Glu Thr Phe Thr Ser Lys Ser Ala Ile Ser Lys Thr Ala
- Glu Ser Ser Gly Gly Pro Ser Thr Ser Arg Val Thr Asp Pro Gln Gly
- Lys Ser Asp Ile Pro Met Asp Leu Phe Asp Phe Tyr Glu Gln Met Asp

Lys Asp Glu Glu Glu Glu Glu Thr Gln Thr Val Ser Phe Glu Val Lys Gln Glu Met Ile Glu Glu Leu Gln Lys Arg Cys Ile His Leu Glu Tyr Pro Leu Leu Ala Glu Tyr Asp Phe Arg Asn Asp Ser Val Asn Pro Asp Ile Asn Ile Asp Leu Lys Pro Thr Ala Val Leu Arg Pro Tyr Gln Glu Lys Ser Leu Arg Lys Met Phe Gly Asn Gly Arg Ala Arg Ser Gly Val Ile Val Leu Pro Cys Gly Ala Gly Lys Ser Leu Val Gly Val Thr Ala Ala Cys Thr Val Arg Lys Arg Cys Leu Val Leu Gly Asn Ser Ala Val Ser Val Glu Gln Trp Lys Ala Gln Phe Lys Met Trp Ser Thr Ile Asp Asp Ser Gln Ile Cys Arg Phe Thr Ser Asp Ala Lys Asp Lys Pro 390 395 Ile Gly Cys Ser Val Ala Ile Ser Thr Tyr Ser Met Leu Gly His Thr Thr Lys Arg Ser Trp Glu Ala Glu Arg Val Met Glu Trp Leu Lys Thr Gln Glu Trp Gly Leu Met Ile Leu Asp Glu Val His Thr Ile Pro Ala Lys Met Phe Arg Arg Val Leu Thr Ile Val Gln Ala His Cys Lys Leu Gly Leu Thr Ala Thr Leu Val Arg Glu Asp Asp Lys Ile Val Asp Leu Asn Phe Leu Ile Gly Pro Lys Leu Tyr Glu Ala Asn Trp Met Glu Leu Gln Asn Asn Gly Tyr Ile Ala Lys Val Gln Cys Ala Glu Val Trp Cys Pro Met Ser Pro Glu Phe Tyr Arg Glu Tyr Val Ala Ile Lys Thr Lys 520 Lys Arg Ile Leu Leu Tyr Thr Met Asn Pro Asn Lys Phe Arg Ala Cys Gln Phe Leu Ile Lys Phe His Glu Arg Arg Asn Asp Lys Ile Ile Val Phe Ala Asp Asn Val Phe Ala Leu Lys Glu Tyr Ala Ile Arg Leu Asn Lys Pro Tyr Ile Tyr Gly Pro Thr Ser Gln Gly Glu Arg Met Gln Ile 580

- Leu Gln Asn Phe Lys His Asn Pro Lys Ile Asn Thr Ile Phe Ile Ser
- Lys Val Gly Asp Thr Ser Phe Asp Leu Pro Glu Ala Asn Val Leu Ile
- Gln Ile Ser Ser His Gly Gly Ser Arg Arg Gln Glu Ala Gln Arg Leu
- Gly Arg Val Leu Arg Ala Lys Lys Gly Met Val Ala Glu Glu Tyr Asn
- Ala Phe Phe Tyr Ser Leu Val Ser Gln Asp Thr Gln Glu Met Ala Tyr
- Ser Thr Lys Arg Gln Arg Phe Leu Val Gln Gly Tyr Ser Phe Lys Val 680
- Ile Thr Lys Leu Ala Gly Met Glu Glu Asp Leu Ala Phe Ser Thr
- Lys Glu Glu Gln Gln Leu Leu Gln Lys Val Leu Ala Ala Thr Asp
- Leu Asp Ala Glu Glu Glu Val Val Ala Gly Glu Phe Gly Ser Arg Ser
- Ser Gln Ala Ser Arg Arg Phe Gly Thr Met Ser Ser Met Ser Gly Ala
- Asp Asp Thr Val Tyr Met Glu Tyr His Ser Ser Arg Ser Lys Ala Pro
- Ser Lys His Val His Pro Leu Phe Lys Arg Phe Arg Lys

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids

 - (B) TYPE: amino acid (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
- (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..15
 - (D) OTHER INFORMATION: /note= "Peptide # 464: XPB peptide that binds wild-type p53 protein."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
 - Leu Gly Leu Thr Ala Thr Leu Val Arg Glu Asp Asp Lys Ile Val 10

- (2) INFORMATION FOR SEQ ID NO:5:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..15
 - (D) OTHER INFORMATION: /note= "Peptide # 479: XPB peptide that does not bind wild-type p53 protein."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Asp Leu Asn Phe Leu Ile Gly Pro Lys Leu Tyr Glu Ala Asn Trp

- (2) INFORMATION FOR SEQ ID NO:6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: Homo sapiens
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide (B) LOCATION: 1..16

 - (D) OTHER INFORMATION: /note= "Peptide # 99: control peptide."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Gly Leu Ser Ala Met Ser Thr Thr Asp Leu Glu Ala Tyr Phe Lys Asp